

## <u>SEQUENCE LISTING</u> (37 C.F.R. §§ 1.821 - 1.825)

(I) GENERAL INFORMATION:

- (i) APPLICANT: ROBERT WEBBER
- (ii) TITLE OF INVENTION: IMMUNOASSAY METHOD EMPLOYING MONOCLONAL ANTIBODY REACTIVE TO HUMAN INOS
- (iii) NUMBER OF SEQUENCES: 126
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: BIELEN, PETERSON & LAMPE
  - (B) STREET: 1990 N. CALIFORNIA BOULEVARD, SUITE 720
  - (C) CITY: WALNUT CREEK
  - (D) STATE: CALIFORNIA
  - (E) COUNTRY: UNITED STATES OF AMERICA
  - (F) ZIP: 94596
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: DISKETTE 3.5 INCH, 1.44 MB FOR FORMATTED
  - (B) COMPUTER: IBM PC COMPATIBLE
  - (C) OPERATING SYSTEM: DOS
  - (D) SOFTWARE: WORDPERFECT 5.1
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: 08/833,506
  - (B) FILING DATE: 7 April 1997
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: 08/634,332
  - (B) FILING DATE: 12 APRIL 1996
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: THEODORE J. BIELEN, JR.
  - (B) REGISTRATION NUMBER: 27,420
  - (C) REFERENCE/DOCKET NUMBER: 12280
- (ix) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: (925) 937-1515
  - (B) TELEFAX: (925) 937-1529
  - (2) INFORMATION FOR SEQ ID NO: 1:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (25-42)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:

## CERTIFICATE OF MAILING

By Signature



- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
- Asn Asn Asn Val Glu Lys Ala Pro Cys Ala Thr Ser Ser

Pro Val Thr Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 2:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: MOUSE iNOS (25-42)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Asn Asn Asn Val Lys Lys Thr Pro Cys Ala Val Leu Ser
5 10
Pro Thr Ile Gln Asp
15

- (2) INFORMATION FOR SEQ ID NO: 3:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: RAT iNOS (25-42)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Asn Asn Asn Val Glu Lys Thr Pro Gly Ala Ile Pro 5 10 Pro Thr Gln Thr Asp 15

- (2) INFORMATION FOR SEQ ID NO: 4:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE

- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (37-54)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Val Gln Asp Asp Leu Gln Ser Pro Thr Tyr His Asn Leu 10

Ser Lys Gln Gln Asn 15

- (2) INFORMATION FOR SEQ ID NO: 5:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (781-798)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10

Gly Pro Thr Pro His 15

- (2) INFORMATION FOR SEQ ID NO: 6:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: MOUSE iNOS (776-792)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- SEQUENCE DESCRIPTION: SEQ ID NO: 6: (xi)

Xaa Ala Gln Gly Ile Leu Glu Arg Val Val Leu Val 5 10

Cys ProThr Pro His 15

(2) INFORMATION FOR SEQ ID NO: 7: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: RAT iNOS (780-794) (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 7: Gln Gly Ile Leu Glu Arg Val Val Asp Xaa Xaa Leu Val 5 10 Xaa Cys Ser Ser Pro 15 (2) INFORMATION FOR SEQ ID NO: 8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (985-1002) (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 8: (xi) Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu 10 5 His Gln His Asp Ser 15 (2) INFORMATION FOR SEQ ID NO: 9: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: MOUSE INOS (978-995) (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION:

SEQUENCE DESCRIPTION: SEQ ID NO: 9:

(xi)

- Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
  5 10
  His Asp Ser Gln His
  15
  - (2) INFORMATION FOR SEQ ID NO: 10:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: RAT iNOS (982-998)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
- Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
  5 10
  His Asp Ser Gln His
- His Asp Ser Gln His
  - (2) INFORMATION FOR SEQ ID NO: 11:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN nNOS (1256-1273)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
- Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Gln
  5
  10

Phe Asp Ile Gln His

- (2) INFORMATION FOR SEQ ID NO: 12:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN eNOS (1017-1031)

- (B) LOCATION:
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
- Gly Ile Ala Pro Phe Arg Gly Phe Trp Gln Glu Arg Leu
  5 10
- His Asp Xaa Xaa Xaa 15
  - (2) INFORMATION FOR SEQ ID NO: 13:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: BOVINE eNOS (1019-1033)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
- Gly Ile Ala Pro Phe Arg Gly Phe Trp Gln Glu Arg Leu
  5 10
- His Asp Xaa Xaa Xaa 15
  - (2) INFORMATION FOR SEQ ID NO: 14:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (1009-1026)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
- Arg Met Thr Leu Val Phe Gly Cys Arg Arg Pro Asp Glu 5
- Asp His Ile Tyr Gln 15
  - (2) INFORMATION FOR SEQ ID NO: 15:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18

- (B) TYPE: AMINO ACID
- (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: RAT iNOS (1006-1023)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

Arg Met Thr Leu Val Phe Gly Cys Arg His Pro Glu Glu 5

Asp His Leu Tyr Gln 15

- (2) INFORMATION FOR SEQ ID NO: 16:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: MOUSE INOS (1002-1019)
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

Arg Met Ser Leu Val Phe Gly Cys Arg His Pro Glu Glu 5

Asp His Leu Tyr Gln 15

- (2) INFORMATION FOR SEQ ID NO: 17:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 16
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: hnNOS  $[2-16, \text{Cys}^{17}]$
  - (B) LOCATION: HUMAN NNOS: AMINO TERMINAL
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

Glu Asp His Met Phe Gly Val Gln Gln Ile Gln Pro Asn
5 10

Val Ile Cys 15

7

- (2) INFORMATION FOR SEQ ID NO: 18:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 24
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: hnNOS  $[Cys^{1410}-1411-1433]$
  - (B) LOCATION: HUMAN NNOS: CARBOXYL TERMINAL
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
- Cys Arg Leu Arg Ser Glu Ser Ile Ala Phe Ile Glu Glu
  5 10
- Ser Lys Lys Asp Thr Asp Glu Val .Phe Ser Ser 15
  - (2) INFORMATION FOR SEQ ID NO: 19:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 20
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: hiNOS  $[2-21, Ser^2]$
  - (B) LOCATION: HUMAN INOS: AMINO TERMINAL
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
- Ala Ser Pro Trp Lys Phe Leu Phe Lys Thr Lys Phe His
  5
- Gln Tyr Ala Met Asn Gly Glu 15 20
  - (2) INFORMATION FOR SEQ ID NO: 20:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: hiNOS [Cys<sup>1136</sup>-1137-1153]
  - (B) LOCATION: HUMAN INOS: CARBOXYL TERMINAL
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

- Cys Lys Lys Asp Arg Val Ala Val Gln Pro Ser Ser Leu
  5 10
  Glu Met Ser Ala Leu
  15
  - (2) INFORMATION FOR SEQ ID NO: 21:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: heNOS [Cap-2-12, Cys<sup>13</sup>]
  - (B) LOCATION: HUMAN ENOS: AMINO TERMINAL WITH CAPROIC ACID ATTACHED
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
- Gly Asn Leu Lys Ser Val Ala Gln Glu Pro Gly Cys
  5 10
  - (2) INFORMATION FOR SEQ ID NO: 22:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: henos  $[2-12, \text{Cys}^{13}]$
  - (B) LOCATION: HUMAN ENOS: AMINO TERMINAL WITHOUT CAPROIC ACID ATTACHED
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
- Gly Asn Leu Lys Ser Val Ala Gln Glu Pro Gly Cys
  5 10
  - (2) INFORMATION FOR SEQ ID NO: 23:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 23
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: heNOS [Cys<sup>1181</sup>-1182-1203]
  - (B) LOCATION: HUMAN ENOS: CARBOXYL TERMINAL
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

Cys Glu Arg Gln Leu Arg Glu Ala Val Pro Trp Ala Phe
5 10

Asp Pro Pro Gly Ser Asp Thr Asn Ser Pro

- (2) INFORMATION FOR SEQ ID NO: 24:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: hiNOS [985-1002]
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:
- Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
  5
  10
- His Asp Ser Gln His
  - (2) INFORMATION FOR SEQ ID NO: 25:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: hiNOS [985-1002]
  - (B) LOCATION:
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:
- Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
  5 10
- His Asp Ser Gln His
  - (2) INFORMATION FOR SEQ ID NO: 26:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR

(ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: hiNOS [37-54] (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 26: (xi) Val Thr Gln Asp Asp His Asn Ser Pro Leu Gln Tyr Leu 5 10 Ser Lys Gln Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 27: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: hiNOS [781-798] (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 27: (xi) Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val 5 10 Gly His Pro Thr Pro 15 (2) INFORMATION FOR SEQ ID NO: 28: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: hiNOS [25-42] (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

Glu Lys Ala Pro Asn Asn Asn Val Ser Ala Thr Ser Ser 10 Pro Val Thr Gln Asp 15

(2) INFORMATION FOR SEQ ID NO: 29: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: hinos [37-54] (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 29: (xi) Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu 10 Ser Lys Gln Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 30: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: hinos [781-798] (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 30: (xi) Pro Ala Leu Gln Gly Ile Leu Glu Arg Val Val Val 5 10 Gly Thr His Pro Pro 15 (2) INFORMATION FOR SEQ ID NO: 31: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: hiNOS [1009-1026] (B) LOCATION: (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

Arq Met Thr Leu Val Phe Gly Ser Arg Arq Pro Asp Glu 5 10 Tyr His Ile Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 32:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: (A3) LOCUS HUMAN iNOS (25-42)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

Ala Asn Asn Asn Val Glu Lys Ala Pro Ser Thr Ser Ser 10 Pro Val Thr Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO: 33:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: MOUSE iNOS (25-42)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

Asn Asn Asn Val Lys Lys Thr Pro Ser Ala Val Leu Ser
5 10
Pro Thr Ile Gln Asp
15

- (2) INFORMATION FOR SEQ ID NO: 34:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: RAT iNOS (25-42)

- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS(D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

Asn Asn Asn Val Glu Lys Thr Pro Gly Ala Ile Pro Ser
5 10
Pro Thr Thr Gln Asp
15

- (2) INFORMATION FOR SEQ ID NO: 35:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (28-42)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35:
- Val Glu Lys Ala Pro Ser Ala Thr Ser Ser Pro Val Thr
  5 10
  Gln Asp
- GIN ASP 15
  - (2) INFORMATION FOR SEQ ID NO: 36:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (31-42)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:
- Ala Pro Ser Ala Thr Ser Ser Pro Val Thr Gln Asp
  5 10
  - (2) INFORMATION FOR SEQ ID NO: 37:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 9
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR

(ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (34-42) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: (xi) SEQ ID NO: 37: Ala Ser Pro Val Thr Gln Asp Thr Ser 5 (2) INFORMATION FOR SEQ ID NO: 38: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-42) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 38: (xi) Ser Pro Val Thr Gln Asp 5 (2) INFORMATION FOR SEQ ID NO: 39: SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (25-39) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39: Asn Asn Asn Val Glu Lys Ala Pro Ser Ala Thr Ser 5 10 Pro Val 15 (2) INFORMATION FOR SEQ ID NO: 40: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID

(D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN iNOS (25-36) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40: Asn Asn Asn Val Glu Lys Ala Pro Ser Ala Thr Ser 5 10 (2) INFORMATION FOR SEQ ID NO: 41: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (25-33) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 41: Asn Asn Asn Val Glu Lys Ala Pro Ser 5 (2) INFORMATION FOR SEQ ID NO: 42: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (25-30) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42: Asn Asn Val Glu Lys (2) INFORMATION FOR SEQ ID NO: 43: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18

(B) TYPE: AMINO ACID(D) TOPOLOGY: LINEAR

(ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: (A4) LOCUS HUMAN INOS (37-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 43: Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu 10 Ser Lys Gln Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 44: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (40-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 44: Thr Gln Asp Asp Leu Gln Tyr His Asn Leu Ser Lys Gln 10 Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 45: (i) **SEQUENCE CHARACTERISTICS:** (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (43-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 45:

(D) OTHER INFORMATION:

Asp Leu Gln Tyr His Asn Leu Ser Lys Gln Gln Asn 5 10

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(2) INFORMATION FOR SEQ ID NO: 46: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (46-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 46: Tyr His Asn Leu Ser Lys Gln Gln Asn 5 (2) INFORMATION FOR SEQ ID NO: 47: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (49-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 47: (xi) Leu Ser Lys Gln Gln Asn (2) INFORMATION FOR SEQ ID NO: 48: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-51) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 48: Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn 5 10 Ser Lys

15

(2) INFORMATION FOR SEQ ID NO: 49: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (37-48) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 49: Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn 5 10 (2) INFORMATION FOR SEQ ID NO: 50: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 50: Ser Pro Val Gln Asp Asp Leu Gln Thr 5 (2) INFORMATION FOR SEQ ID NO: 51: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-42) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 51:

Ser Pro Val

Thr

Gln

5

Asp

(2) INFORMATION FOR SEQ ID NO: 52: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: (F6) LOCUS HUMAN INOS (781-798) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 52: (xi) Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10 Pro His Gly Pro Thr 15 (2) INFORMATION FOR SEQ ID NO: 53: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN eNOS (806-824) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 53: (xi) Glu Ala Leu Leu Ser Pro Gly Leu Val Arg Val Glu Asp 10 Thr Pro Pro Ala Pro Glu 15 (2) INFORMATION FOR SEQ ID NO: 54: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (784-798) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 54: (xi)

Gln Gly Ile Leu Glu Arg Val Val Asp Gly Pro Thr 10 Pro His 15 (2) INFORMATION FOR SEQ ID NO: 55: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (787-798) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 55: (xi) Ile Leu Glu Arg Val Val Asp Gly Pro Thr Pro His 5 10 (2) INFORMATION FOR SEQ ID NO: 56: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (790-798) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 56: (xi) Arg Val Val Asp Gly Pro Thr Pro His 5 (2) INFORMATION FOR SEQ ID NO: 57: SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (793-798) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION:

SEQUENCE DESCRIPTION: SEQ ID NO: 57:

(xi)

Asp Gly Pro Thr Pro His
5

- (2) INFORMATION FOR SEQ ID NO: 58:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 14
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (781-794)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 58:

Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp 5 10 Gly

- (2) INFORMATION FOR SEQ ID NO: 59:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (781-792)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 59:

Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val 5

- (2) INFORMATION FOR SEQ ID NO: 60:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 9
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (781-789)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 60:

Pro Ala Leu Val Gln Gly Ile Leu Glu 5

- (2) INFORMATION FOR SEQ ID NO: 61:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 6
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (781-786)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 61:

Pro Ala Leu Val Gln Gly 5

- (2) INFORMATION FOR SEQ ID NO: 62:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: (G11) LOCUS HUMAN iNOS (985-1002)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 62:
- Gly Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu
  5 10
- His Asp Ser Gln His
  - (2) INFORMATION FOR SEQ ID NO: 63:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN nNOS (1256-1273)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 63:

Gly Ile Ala Pro Phe Arg Ser Phe Trp Gln Gln Arg Gln
5 10
Phe Asp Ile Gln His
15

- (2) INFORMATION FOR SEQ ID NO: 64:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN eNOS (1017-1031)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 64:
- Gly Ile Ala Pro Phe Arg Gly Phe Trp Gln Glu Arg Leu
  5 10

His Asp 15

- (2) INFORMATION FOR SEQ ID NO: 65:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (988-1002)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 65:
- Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu His Asp Ser 5
- Gln His
  - (2) INFORMATION FOR SEQ ID NO: 66:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (991-1002)

- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEO ID NO: 66:

Ser Phe Trp Gln Gln Arg Leu His Asp Ser Gln His 5 10

- (2) INFORMATION FOR SEQ ID NO: 67:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 9
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- MOLECULE TYPE: PEPTIDE (ii)
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (994-1002)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- SEQUENCE DESCRIPTION: SEQ ID NO: 67: (xi)
- Gln Gln Arg Leu His Asp Ser Gln His 5
  - (2) INFORMATION FOR SEQ ID NO: 68:
- (i)SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 5
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- MOLECULE TYPE: PEPTIDE (ii)
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (997-1002)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 68:

His Asp Ser Gln His 5

- (2) INFORMATION FOR SEQ ID NO: 69:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (985-998)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 69: Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu Gly 5 10 His Asp 15 (2) INFORMATION FOR SEQ ID NO: 70: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (985-996) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 70: (xi) Gly Ile Val Gln Gln Arg Pro Phe Arg Ser Phe Trp 5 10 (2) INFORMATION FOR SEQ ID NO: 71: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (985-993) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 71: (xi) Gly Ile Val Phe Arg Ser Phe Trp Pro 5 (2) INFORMATION FOR SEQ ID NO: 72: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (985-990) 26

- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 72:
- Gly Ile Val Pro Phe Arg
  - (2) INFORMATION FOR SEQ ID NO: 73:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: (H1) LOCUS HUMAN INOS (1009-1026)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 73:
- Arg Met Thr Leu Val Phe Gly Ser Arg Arg Pro Asp Glu
  5 10
  Asp His Ile Tyr Gln
  - (2) INFORMATION FOR SEQ ID NO: 74:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 17
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:

15

- (A) NAME/KEY: HUMAN eNOS (1041-1057)
- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 74:
- Met Thr Leu Val Phe Gly Ser Arg Ser Ser Gln Leu Asp
  5 10
  His Leu Tyr Arg

His Leu Tyr Arg

- (2) INFORMATION FOR SEQ ID NO: 75:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 17
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR

- (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN nNOS (1281-1297) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 75: (xi) Met Val Phe Gly Ser Arg Gln Ser Lys Ile Asp Leu Val 5 10 His Ile Tyr Arg 15 (2) INFORMATION FOR SEQ ID NO: 76: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (1012-1026) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 76: (xi) Phe Gly Val Ser Arg Arg Pro Asp Glu Asp His Ile 10
  - (2) INFORMATION FOR SEQ ID NO: 77:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:

Gln 15

Tyr

- (A) NAME/KEY: HUMAN INOS (1015-1026)
- (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
- (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
- (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 77:
- Gly Ser Arg Arg Pro Asp Glu Asp His Ile Tyr Gln
  5 10

(2) INFORMATION FOR SEQ ID NO: 78: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN INOS (1018-1026) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 78: (xi) Glu Asp His Ile Tyr Gln Arg Pro Asp 5 (2) INFORMATION FOR SEQ ID NO: 79: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) **FEATURE:** (A) NAME/KEY: HUMAN iNOS (1021-1026) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 79: (xi) Tyr Glu Asp His Ile Gln 5 (2) INFORMATION FOR SEQ ID NO: 80: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (1009-1023) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 80: Arg Met Thr Leu Val Phe Gly Ser Arq Pro Arq Asp 5 10 Asp His 15

(2) INFORMATION FOR SEQ ID NO: 81: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (1009-1020) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 81: Arg Met Thr Leu Val Phe Gly Ser Arg Arg Pro 5 10 (2) INFORMATION FOR SEQ ID NO: 82: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (1009-1017) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 82: (xi) Arg Met Thr Leu Val Phe Gly Ser Arg 5 (2) INFORMATION FOR SEQ ID NO: 83: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (1009-1014) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 83: Arq Met Thr Val Phe Leu 5

(2) INFORMATION FOR SEQ ID NO: 84: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: TRUNCATED HUMAN INOS (40-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 84: (xi) Thr Gln Asp Asp Leu Gln Tyr His Asn Leu Ser Lys 5 10 (2) INFORMATION FOR SEQ ID NO: 85: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: TRUNCATED HUMAN INOS (784-798) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 85: (xi) Val Gln Gly Ile Leu Glu Arg Val Val 5 (2) INFORMATION FOR SEQ ID NO: 86: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 86: (xi) Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu Ser Pro

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5

Asn

Gln

Ser

Lys

15

Gln

(2) INFORMATION FOR SEQ ID NO: 87: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (41-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 87: (xi) Gln Asp Asp Leu Gln 5 (2) INFORMATION FOR SEQ ID NO: 88: (i)SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE **FEATURE:** (ix) (A) NAME/KEY: HUMAN iNOS (40-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 88: (xi) Leu Gln Thr Gln Asp Asp 5 (2) INFORMATION FOR SEQ ID NO: 89: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (39-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 89: (xi) Val Thr Gln Asp Leu Gln Asp

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(2) INFORMATION FOR SEQ ID NO: 90: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (38-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 90: Pro Val Thr Gln Asp Asp Leu Gln 5 (2) INFORMATION FOR SEQ ID NO: 91: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-45) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 91: Ser Pro Val Thr Gln Asp Asp Leu Gln (2) INFORMATION FOR SEQ ID NO: 92: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (40-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 92: Thr Gln Asp Leu Asp 5

(2) INFORMATION FOR SEQ ID NO: 93: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (39-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 93: Thr Gln Asp Val Asp Leu 5 (2) INFORMATION FOR SEQ ID NO: 94: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (38-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 94: (xi) Pro Val Thr Gln Asp Asp Leu (2) INFORMATION FOR SEQ ID NO: 95: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 95: Ser Pro Val Thr Gln Asp Asp Leu

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(2) INFORMATION FOR SEQ ID NO: 96: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (36-44) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 96: Ser Ser Pro Val Thr Gln Asp Asp Leu 5 (2) INFORMATION FOR SEQ ID NO: 97: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (39-43) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 97: (xi) Val Thr Gln Asp Asp 5 (2) INFORMATION FOR SEQ ID NO: 98: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (38-43) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 98: (xi) Thr Pro Val Asp Gln Asp

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- (2) INFORMATION FOR SEQ ID NO: 99: (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 7
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (37-43)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 99:

Ser Pro Val Thr Gln Asp Asp 5

- (2) INFORMATION FOR SEQ ID NO: 100:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 8
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (36-43)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 100:

Ser Ser Pro Val Thr Gln Asp Asp 5

- (2) INFORMATION FOR SEQ ID NO: 101:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 9
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (35-43)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 101:

Thr Ser Ser Pro Val Thr Gln Asp Asp 5

(2) INFORMATION FOR SEQ ID NO: 102: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (37-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: (xi) **SEQ ID NO: 102:** Val Tyr His Asn Leu Ser Pro Thr Gln Asp Asp Leu Gln 5 10 Ser Lys Gln Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 103: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (40-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 103: (xi) Gln Asp Thr Asp Leu Gln Tyr His Asn Leu Ser Lys Gln 5 10 Gln Asn 15 (2) INFORMATION FOR SEQ ID NO: 104: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (43-54) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION:

SEQUENCE DESCRIPTION: SEQ ID NO: 104:

(xi)

Asp Leu Gln Tyr His Asn Leu Ser Lys Gln Gln Asn 5 10

- (2) INFORMATION FOR SEQ ID NO: 105:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 9
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (46-54)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 105:

Tyr His Asn Ser Lys Gln Gln Asn Leu 5

- (2) INFORMATION FOR SEQ ID NO: 106:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 6
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- MOLECULE TYPE: PEPTIDE (ii)
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (49-54)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 106:

Leu Ser Lys Gln Gln Asn 5

- (2) INFORMATION FOR SEQ ID NO: 107:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (37-51)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 107:

Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn Leu 5 10 Ser Lys 15

- (2) INFORMATION FOR SEQ ID NO: 108:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 12
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (37-48)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 108:
- Ser Pro Val Thr Gln Asp Asp Leu Gln Tyr His Asn 5
  - (2) INFORMATION FOR SEQ ID NO: 109:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 9
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (37-45)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 109:
- Ser Pro Val Thr Gln Asp Asp Leu Gln 5
  - (2) INFORMATION FOR SEQ ID NO: 110:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 6
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (37-42)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 110:

Ser Pro Val Thr Gln Asp 5

- (2) INFORMATION FOR SEQ ID NO: 111:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 10
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (35-44)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

Thr Ser Ser Pro Val Thr Gln Asp Asp Leu 5 10

- (2) INFORMATION FOR SEQ ID NO: 112:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (781-798)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 112:
- Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp
- Gly Pro Thr Pro His
  - (2) INFORMATION FOR SEQ ID NO: 113:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 5
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (788-792)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 113: Leu Glu Arg Val Val 5 (2) INFORMATION FOR SEQ ID NO: 114: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (787-792) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 114: (xi) Ile Leu Glu Arq Val Val 5 (2) INFORMATION FOR SEQ ID NO: 115: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN INOS (786-792) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 115: (xi) Gly Ile Leu Glu Arg Val Val 5 (2) INFORMATION FOR SEQ ID NO: 116: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN INOS (785-792) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE

(C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(D) OTHER INFORMATION:

SEQUENCE DESCRIPTION: SEQ ID NO: 116: Gln Gly Ile Leu Glu Arg Val Val 5 (2) INFORMATION FOR SEQ ID NO: 117: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (784-792) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 117: Val Gln Gly Ile Leu Glu Arg Val Val 5 (2) INFORMATION FOR SEQ ID NO: 118: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (787-791) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 118: (xi) Ile Leu Glu Arg Val 5 (2) INFORMATION FOR SEQ ID NO: 119: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (786-791) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 119: Gly Ile Leu Glu Arg Val 5 (2) INFORMATION FOR SEQ ID NO: 120: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE FEATURE: (ix) (A) NAME/KEY: HUMAN iNOS (785-791) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 120: (xi) Gln Gly Ile Leu Glu Arg Val 5 (2) INFORMATION FOR SEQ ID NO: 121: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR MOLECULE TYPE: PEPTIDE (ii) (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (784-791) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS (D) OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 121: (xi) Gln Gly Ile Leu Glu Arg Val Val (2) INFORMATION FOR SEQ ID NO: 122: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 (B) TYPE: AMINO ACID (D) TOPOLOGY: LINEAR (ii) MOLECULE TYPE: PEPTIDE (ix) FEATURE: (A) NAME/KEY: HUMAN iNOS (783-791) (B) LOCATION: CARBOXY TERMINAL WITH AMIDE (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS

(D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 122:

Leu Val Gln Gly Ile Leu Glu Arg Val

- (2) INFORMATION FOR SEQ ID NO: 123:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 5
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (786-790)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 123:
- Gly Ile Leu Glu Arg
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  - (2) INFORMATION FOR SEQ ID NO: 124:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 6
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (785-790)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 124:
- Gln Gly Ile Leu Glu Arg
  5
  - (2) INFORMATION FOR SEQ ID NO: 125:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 7
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (784-790)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 125:

Val Gln Gly Ile Leu Glu Arg
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- (2) INFORMATION FOR SEQ ID NO: 126:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 8
  - (B) TYPE: AMINO ACID
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: PEPTIDE
- (ix) FEATURE:
  - (A) NAME/KEY: HUMAN iNOS (783-790)
  - (B) LOCATION: CARBOXY TERMINAL WITH AMIDE
  - (C) IDENTIFICATION METHOD: AMINO ACID ANALYSIS
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 126:

Leu Val Gln Gly Ile Leu Glu Arg